

SEQUENCE LISTING

<110> COMMISSARIAT A L'ENERGIE ATOMIQUE
UNIVERSITE PIERRE ET MARIE CURIE (PARIS VI)

<120> LABELLED PEPTIDES HAVING AN AFFINITY FOR A PHOSPHOLIPID
AND USES

<130> B14023EE

<140> PCT/FR03/02027
<141> 2003-06-30

<150> FR N°02 08204
<151> 2002-07-01

<160> 14

<170> PatentIn Ver. 2.1

<210> 1
<211> 75
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: sequence
derived from a human annexin

<400> 1
Gly Phe Asp Glu Arg Ala Asp Val Glu Thr Leu Arg Lys Ala Met Lys
1 5 10 15
Gly Leu Gly Thr Asp Glu Glu Ser Ile Leu Thr Leu Leu Thr Ser Arg
20 25 30
Ser Asn Ala Gln Arg Gln Glu Ile Ser Ala Ala Tyr Lys Thr Leu Phe
35 40 45
Gly Arg Asp Leu Leu Asp Asp Leu Lys Ser Glu Leu Thr Gly Lys Phe
50 55 60
Glu Lys Leu Val Val Ala Leu Leu Lys Pro Ser
65 70 75

<210> 2
<211> 75
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: sequence
derived from a human annexin

<400> 2
Asn Phe Asp Ala Glu Arg Asp Ala Leu Asn Ile Arg Lys Ala Ile Lys
1 5 10 15
Gly Met Gly Val Asp Glu Asp Thr Ile Val Asn Ile Leu Thr Asn Arg
20 25 30

<220>
<223> Description of Artificial Sequence: sequence
derived from a human annexin

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<210> 4
<211> 75
<212> PRT
<213> Artificial Sequence
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<220>
<223> Description of Artificial Sequence: sequence
derived from a human annexin

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<400> 4
Gly Phe Asn Ala Met Glu Asp Ala Gln Thr Leu Arg Lys Ala Met Lys
  1          5          10          15
Gly Leu Gly Thr Asp Glu Asp Ala Ile Ile Ser Val Leu Ala Tyr Arg
          20          25          30
Asn Thr Ala Gln Arg Gln Glu Ile Arg Thr Ala Tyr Lys Ser Thr Ile
          35          40          45
Gly Arg Asp Leu Ile Asp Asp Leu Lys Ser Glu Leu Ser Gly Asn Phe
          50          55          60
Glu Arg Val Ile Val Gly Met Met Thr Pro Ser
  65          70          75

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<210> 5
 <211> 75
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: sequence
 derived from a human annexin

<400> 5
 Gly Phe Asp Pro Asn Gln Asp Ala Glu Ala Leu Arg Thr Ala Met Lys
 1 5 10 15
 Gly Phe Gly Ser Asp Glu Glu Ala Ile Leu Asp Ile Ile Thr Ser Arg
 20 25 30
 Ser Asn Arg Gln Arg Gln Glu Val Cys Gln Ser Tyr Lys Ser Leu Tyr
 35 40 45
 Gly Arg Asp Leu Ile Ala Asp Leu Lys Ser Glu Leu Thr Gly Lys Phe
 50 55 60
 Glu Arg Leu Ile Val Gly Leu Met Arg Pro Ser
 65 70 75

<210> 6
 <211> 75
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: sequence
 derived from a human annexin

<400> 6
 Gly Phe Asn Pro Asp Ala Asp Ala Lys Ala Leu Arg Lys Ala Met Lys
 1 5 10 15
 Gly Leu Gly Thr Asp Glu Asp Thr Ile Ile Asp Ile Ile Thr His Arg
 20 25 30
 Ser Asn Val Gln Arg Gln Gln Ile Arg Gln Thr Phe Lys Ser His Phe
 35 40 45
 Gly Arg Asp Leu Met Thr Asp Leu Lys Ser Glu Ile Ser Gly Asp Leu
 50 55 60
 Glu Arg Leu Ile Leu Gly Leu Met Met Pro Ser
 65 70 75

<210> 7
 <211> 75
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: sequence

derived from a human annexin

<400> 7

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Pro Gly Asp Ala Ile Arg Asp Ala Glu Ile Leu Arg Lys Ala Met Lys
 1           5           10           15
Gly Phe Gly Thr Asp Glu Gln Ala Ile Val Asp Val Val Ala Asn Arg
          20           25           30
Ser Asn Asp Gln Arg Gln Lys Ile Lys Ala Ala Phe Lys Thr Ser Tyr
          35           40           45
Gly Arg Asp Leu Ile Lys Asp Leu Lys Ser Glu Leu Ser Gly Asn Met
          50           55           60
Glu Arg Leu Ile Leu Ala Leu Phe Met Pro Ser
          65           70           75

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<210> 8

<211> 75

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: sequence
derived from a human annexin

<400> 8

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His Phe Asn Pro Asp Pro Asp Val Glu Thr Leu Arg Lys Ala Met Lys
 1           5           10           15
Gly Ile Gly Thr Asn Glu Gln Ala Ile Ile Asp Val Leu Thr Lys Arg
          20           25           30
Ser Asn Thr Gln Arg Gln Thr Ile Ala Lys Ser Phe Lys Ala Gln Phe
          35           40           45
Gly Arg Asp Leu Thr Glu Asp Leu Lys Ser Glu Leu Ser Gly Lys Leu
          50           55           60
Glu Arg Leu Ile Val Ala Leu Met Tyr Pro Ser
          65           70           75

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<210> 9

<211> 75

<212> PRT

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<223> Description of Artificial Sequence: sequence
derived from a human annexin

<400> 9

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Gly Phe Asp Pro Leu Arg Asp Ala Glu Val Leu Arg Lys Ala Met Lys
 1           5           10           15
Gly Phe Gly Thr Asp Glu Gln Ala Ile Ile Asp Cys Leu Gly Ser Arg
          20           25           30

```

Ser Asn Lys Gln Arg Gln Gln Ile Leu Leu Ser Phe Lys Thr Ala Tyr
 35 40 45

Gly Arg Asp Leu Ile Lys Asp Leu Lys Ser Glu Leu Ser Gly Asn Phe
 50 55 60

Glu Lys Thr Ile Leu Ala Leu Met Lys Thr Ser
 65 70 75

<210> 10

<211> 75

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: sequence
 derived from a human annexin

<400> 10

Gly Phe Asp Val Asp Arg Asp Ala Lys Lys Leu Arg Lys Ala Met Lys
 1 5 10 15

Gly Met Gly Thr Asn Glu Ala Ala Ile Ile Glu Ile Leu Ser Gly Arg
 20 25 30

Thr Ser Asp Glu Arg Gln Gln Ile Lys Gln Lys Tyr Lys Ala Thr Tyr
 35 40 45

Gly Arg Glu Leu Glu Glu Asp Leu Lys Ser Glu Leu Ser Gly Asn Phe
 50 55 60

Glu Lys Thr Ala Leu Ala Leu Leu Asp Arg Ser
 65 70 75

<210> 11

<211> 79

<212> PRT

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<223> Description of Artificial Sequence: sequence
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<223> Xaa of position 22 is Leu, Met or Trp

<220>

<223> Xaa of position 34 is Thr or Lys

<220>

<223> Xaa of position 45 is Ser or Lys

<220>

<223> Xaa of position 48 is Phe or Tyr

<220>

<223> Xaa of position 50 is Thr or Glu

<220>

<223> Xaa of position 63 is Glu or Lys

<220>

<223> Xaa of position 69 is Glu or Lys

<220>

<223> Xaa of position 71 is Glu or Leu

<400> 11

Gly Ser Gly Cys Gly Phe Asp Glu Arg Ala Asp Val Glu Thr Leu Arg
1 5 10 15

Lys Ala Met Lys Gly Xaa Gly Thr Asp Glu Glu Ser Ile Leu Thr Leu
20 25 30

Leu Xaa Ser Arg Ser Asn Ala Gln Arg Gln Glu Ile Xaa Ala Ala Xaa
35 40 45

Lys Xaa Leu Phe Gly Arg Asp Leu Leu Asp Asp Leu Lys Ser Xaa Leu
50 55 60

Thr Gly Lys Phe Xaa Lys Xaa Val Val Ala Leu Leu Lys Pro Ser
65 70 75

<210> 12

<211> 78

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: sequence
derived from a human annexin

<400> 12

Gly Ser Pro Gly Phe Asp Glu Arg Ala Asp Val Glu Thr Leu Arg Lys
1 5 10 15

Ala Met Lys Gly Leu Gly Thr Asp Glu Glu Ser Ile Leu Thr Leu Leu
20 25 30

Thr Ser Arg Ser Asn Ala Gln Arg Gln Glu Ile Ser Ala Ala Tyr Lys
35 40 45

Thr Leu Phe Gly Arg Asp Leu Leu Asp Asp Leu Lys Ser Glu Leu Thr
50 55 60

Gly Lys Phe Glu Lys Leu Val Val Ala Leu Leu Lys Pro Ser
65 70 75

<210> 13

<211> 83

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: sequence
derived from a human annexin

<220>
 <223> Xaa of position 25 is Leu, Met or Trp

<220>
 <223> Xaa of position 37 is Thr or Lys

<220>
 <223> Xaa of position 48 is Ser or Lys

<220>
 <223> Xaa of position 51 is Phe or Tyr

<220>
 <223> Xaa of position 53 is Thr or Glu

<220>
 <223> Xaa of position 66 is Glu or Lys

<220>
 <223> Xaa of position 72 is Glu or Lys

<220>
 <223> Xaa of position 74 is Glu or Leu

<400> 13
 Gly Ser Glu Cys Asp Phe Pro Gly Phe Asp Glu Arg Ala Asp Val Glu
 1 5 10 15
 Thr Leu Arg Lys Ala Met Lys Gly Xaa Gly Thr Asp Glu Glu Ser Ile
 20 25 30
 Leu Thr Leu Leu Xaa Ser Arg Ser Asn Ala Gln Arg Gln Glu Ile Xaa
 35 40 45
 Ala Ala Xaa Lys Xaa Leu Phe Gly Arg Asp Leu Leu Asp Asp Leu Lys
 50 55 60
 Ser Xaa Leu Thr Gly Lys Phe Xaa Lys Xaa Val Val Ala Leu Leu Lys
 65 70 75 80
 Pro Ser Arg

<210> 14
 <211> 87
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: sequence
 derived from a human annexin

<220>
 <223> Xaa of position 29 is Leu, Met or Trp

<220>
 <223> Xaa of position 41 is Tyr or Lys

<220>

<223> Xaa of position 52 is Ser or Lys

<220>

<223> Xaa of position 55 is Phe or Tyr

<220>

<223> Xaa of position 57 is Thr or Glu

<220>

<223> Xaa of position 70 is Glu or Lys

<220>

<223> Xaa of position 76 is Glu or Lys

<220>

<223> Xaa of position 78 is Glu or Leu

<400> 14

Gly	Ser	Gly	Cys	Gly	Thr	Glu	Thr	Asp	Phe	Pro	Gly	Phe	Asp	Glu	Arg
1				5					10					15	

Ala	Asp	Val	Glu	Thr	Leu	Arg	Lys	Ala	Met	Lys	Gly	Xaa	Gly	Thr	Asp
			20					25					30		

Glu	Glu	Ser	Ile	Leu	Thr	Leu	Leu	Xaa	Ser	Arg	Ser	Asn	Ala	Gln	Arg
		35					40					45			

Gln	Glu	Ile	Xaa	Ala	Ala	Xaa	Lys	Xaa	Leu	Phe	Gly	Arg	Asp	Leu	Leu
	50					55					60				

Asp	Asp	Leu	Lys	Ser	Xaa	Leu	Thr	Gly	Lys	Phe	Xaa	Lys	Xaa	Val	Val
65					70					75					80

Ala	Leu	Leu	Lys	Pro	Ser	Arg
					85	